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January 2, 2006, 06:28:24; Search time 1956 Seconds (without alignments) 10404.366 Million cell updates/sec
                                                                                                                                                                                                                                                                                                  1 gaggaatcgttctgcaaatc......ctgtacttttcatggattag 2461
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cgn2_6/ptodata/1/pubpna/US10A_PUBCOMB.seq:
cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:
cgn2_6/ptodata/1/pubpna/US10C_PUBCOMB.seq:
cgn2_6/ptodata/1/pubpna/US10D_PUBCOMB.seq:
cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:
/cgn2_6/ptodata/1/pubpna/US10B_PUBCOMB.seq:
/cgn2_6/ptodata/1/pubpna/US11_PUBCOMB.seq:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ptodata/1/pubpna/US08_PUBCOMB.seq:
/ptodata/1/pubpna/US09A_PUBCOMB.seq
GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                    9793542 seqs, 4134689005 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Published_Applications_NA_Main:*
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Maximum Match 100%
Listing first 45 summaries
                                                                                                    - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Perfect score:
Sequence:
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                                                                                                                                                      Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

Result		* Query			SUMMAKIES	ť
Š	Score	Match	Match Length DB	8	OI	Description
-	2433.4	98.9	3097	s	US-10-079-625-1	Sequence 1, Appli
7	2433.4	98.9	3656	'n	US-10-226-579-1	Sequence 1, Appli
m	2433.4	98.9	3854	s	US-10-079-625-42	Sequence 42, Appl
4	2390.4	97.1	2679	Ŋ	US-10-278-733-14	14,
S	2364.6	96.1	2868	0	US-08-779-457-8	Sequence 8, Appli
9	2364.6	96.1	2868	'n	US-10-214-802-8	œ
7	2364.6	96.1	2868	œ	US-10-921-710-8	æ
œ	2148.8	87.3	3489	ហ	US-10-278-733-18	18,
6	1692.2	68.8	3102	N	US-08-779-457-6	6, A
2	1692.2	68.8	3102	ß	US-10-214-802-6	9
דו	1692.2	68.8	3102	œ	US-10-921-710-6	6,
17	1692.2	68.8	3800	Ŋ	US-10-226-579-3	
13	1692.2	68.8	3800	7	US-10-641-643-885	88
14	1692.2	68.8	3800	σ	US-10-893-315-12	
15	1692.2	68.8	4102	7	US-08-779-457-1	Sequence 1, Appli
16	1692.2	68.8	4102	Ŋ	US-10-214-802-1	ų
11	1692.2	68.8	4102	00	US-10-921-710-1	Sequence 1, Appli
18	1691	68.7	3004	7	US-08-779-457-5	Sequence 5, Appli
19	1691	68.7	3004	Ŋ	US-10-214-802-5	Sequence 5, Appli
50	1691	68.7	3004	60	US-10-921-710-5	Sequence 5, Appli
21	1690.8	68.7	2991	Ŋ	US-10-095-929-1	Sequence 1, Appli
22	1690.8	68.7	2991	10	US-11-026-133-1	Sequence 1, Appli
23	1689	9.89	3784	0	US-10-893-315-26	, 9

Sequence 9, Appli	Sequence 3, Appli	Sequence 1, Appli	Sequence 12, Appl	Sequence 1, Appli	Sequence 9, Appli	'n	Sequence 7, Appli	13,	Sequence 5, Appli	11,	Sequence 93, Appl	ų,	Sequence 8348, Ap	Sequence 7, Appli	Sequence 8799, Ap	Sequence 13143, A	Sequence 148, App	Sequence 134, App	Sequence 496, App	Sequence 861, App	Sequence 26843, A
US-09-116-676-9	US-10-079-625-3	US-10-245-616-1	US-10-014-156-12	US-10-373-624A-1	US-10-774-721-9	US-10-373-624A-3	US-10-373-624A-7	US-10-774-721-13	US-10-373-624A-5	US-10-774-721-11	US-10-116-802-93	US-10-803-459C-1	US-09-918-995-8348	US-10-803-459C-7	US-09-918-995-8799	US-10-029-386-13143	US-10-893-315-148	US-10-893-315-134	US-10-893-315-496	US-10-893-315-861	US-10-029-386-26843
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9.6	9.8	8.4	68.3	68.1	1.8	9.6	9.	5.6	9.6	9.6	62.6	9.5	1.1	6.0	6.0	9.6	9.6	9.8	9.6	9.8	8.3
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2.8	7.4	1682.8	680.4	1676	1676	1613.8	3.8	3.8	613.8	613.8	1540	480.4	274	267.8	252.4	212.4	212.4	212.4	212	212	205.4
1688	1687	168	168	-	-	161	1613	1613	161	161	-	48		26	25	21	21	21			20
24	25	56	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

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CITY:
STRIE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUT
RESULT 1
US-10-079-625-1
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us-08-783-734d-9.rni

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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Sequence 6, A
Sequence 6, A
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Sequence 1,
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GenCore version 5.1.6
(c) 1993 - 2006 Compugen Ltd.
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US-09-094-410-1

US-08-593-153A-1

US-08-593-153A-1

US-08-638-524B-1

US-08-638-524B-1

US-08-638-524B-4

US-09-094-54A-42

US-09-094-410-42

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US-08-094-410-42

US-08-094-410-42

US-08-638-524B-42

US-08-638-524B-42

US-08-638-524B-42

US-08-827-962-17

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US-08-837-635-5

US-08-833-635-7
                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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US-09-023-655-885
US-09-059-4558-3
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US-08-640-389A-1
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4102
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GENERAL INFORMATION:
APPLICANT: Tartaglia, Louis A.
APPLICANT: Tepper, Robert I.
APPLICANT: Culpepper, Janice A.
TITLE OF INVENTION: METHODS OF IDENTIFYING COMPOUNDS THAT
TITLE OF INVENTION: MODULATE BODY WEIGHT USING THE OB RECEPTOR
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US

ZIP: 02110-2804
COMPUTER READBLE FORM:
WEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: PARESE for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/599,455B
FILING DATE: 22-JAN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/593,153
APPLICATION NUMBER: 08/593,153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07334/017001
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PILING DATE: 18-05-195
PILING DATE: 11-DEC-1995
APPLICATION NUMBER: 08/50,445
APPLICATION NUMBER: 08/56,485
PILING DATE: 08-DEC-1995
APPLICATION NUMBER: 08/56,622
FILING DATE: 04-DEC-1995
APPLICATION NUMBER: 08/56,63
PILING DATE: 27-NOV-1995
ATORNEY/AGENT INFORMATION: NAME: MAKE: MAKE: MAKE: 37-NOV-1995
ATORNEY/AGENT INFORMATION: NAME: NAME: 35,283
                                                                                                                                                                                                                                                                                                                           B: Fish & Richardson, P.C.
225 Franklin Street
RESULT 1
18-08-599-455B-1
Sequence 1, Application US/08599455B
Patent No. 5972621
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-8906
TELEFAX: 617-542-8906
                                                                                                                                                                                                                                                                                                                                                                                         CITY: Boston
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Appl

Sequence

Sequence

Sequence 35377, Sequence Sequence Sequence

Sequence

Sequence

Sequence Sequence Sequence

Sequence 40762, Sequence 13421, Sequence 35776,

Sequence Sequence Sequence

Sequence

OM nucleic

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Run

Sequence:

Searched:

Database

Result

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ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION TUMBER: US/11/192,219
FILING DATE: 27-011-2005
CLASSIFICATION: 424
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US-10-750-185-33603
US-10-750-185-55703
US-10-995-561-459
US-10-750-185-57835
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US-10-750-185-377620
US-10-750-185-377620
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; Publication No. US20050272656A1
GENERAL INPORMATION:
APPLICANT: Matthews, William
ITILE OF INVENTION: USES FOR WSX LIGANDS
; CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/667,197
FILING DATE: 20-Jun-1996
APPLICATION NUMBER: 08/585005
FILING DATE: 08-Jan-96
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: P-40,378
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALIGNMENTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 415/225-1994
TELEFAX: 415/952-988I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
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Sequence 4354, Ap
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                                                                                                                  January 2, 2006, 06:35:53 ; Search time 502 Seconds (without alignments) 2569.973 Million cell updates/sec
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Sequence 38349,
Sequence 27952,
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Sequence 178,
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2461
1 gaggaatcgttctgcaaatc......ctgtactttcatggattag 2461
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Sequence 1
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Sequence 2
Sequence 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO6_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO7_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO1_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO1_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO1_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/USO1_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USO1_NEW_PUB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-536-320-1

US-10-955-561-179

US-10-995-561-179

US-10-793-626-3727

US-10-793-626-4354

US-10-793-626-4354

US-11-117-187-186

US-11-17-187-186

US-11-17-187-186

US-10-750-185-52715

US-10-750-185-24691

US-10-750-185-24691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -10-750-185-38349
-10-750-185-27952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0-185-55839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -10-750-185-46273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1-826A-393
                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                          4172979 segs, 262114271 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Applications NA New:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                               - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                    Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                         Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query
Match Length DB
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135019
67088
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3085
3170
3864
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Published
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Post-processing:
                                                                                                                                                                                                  Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1691
1682.8
```

```
GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
```

OM protein - protein search, using sw model

January 2, 2006, 13:46:09; Search time 8.05474 Seconds (without alignments) 748.439 Million cell updates/sec Run on:

US-08-783-734D-10 4321 1 MMCQKFYVVILHWBFLYVIA......IPSNVKKFYIHGMCTVLFMD 805 Title: Perfect score: Sequence:

**BLOSUM62** Scoring table:

57103 segs, 7488799 residues Gapop 10.0 , Gapext 0.5 Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Total number of hits satisfying chosen parameters:

Post-processing: Minimum Match 08
Maximum Match 1008
Listing first 45 summaries

cgn2\_6/ptodata/2/pubpaa/US06\_NEW\_PUB.pep: /cgn2\_6/ptodata/2/pubpaa/US07\_NEW\_PUB.pep: /cgn2\_6/ptodata/2/pubpaa/PCT\_NEW\_PUB.pep:\* /cgn2\_6/ptodata/2/pubpaa/US09\_NEW\_PUB.pep:\* /cgn2\_6/ptodata/2/pubpaa/US10\_NEW\_PUB.pep:\* /cgn2\_6/ptodata/2/pubpaa/US10\_NEW\_PUB.pep:\* /cgn2\_6/ptodata/2/pubpaa/US10\_NEW\_PUB.pep:\* Published Applications AA New:\*

1: /cgn2\_6/ptodata/2/pubpaa/USO8\_NEW |

2: /cgn2\_6/ptodata/2/pubpaa/USO6\_NEW |

3: /cgn2\_6/ptodata/2/pubpaa/USO7\_NEW |

4: /cgn2\_6/ptodata/2/pubpaa/PCT\_NEW | Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

		40			SOMETES	
Result No.	Score	Query	Query Match Length DB	æ	OI De	Description
-	4031	93.3	783	-	US-11-192-219-7	Sequence 7, Appli
~	3340	77.3	968	,	US-11-192-219-3	m
m	3340	77.3	923	7	4	Sequence 4, Appli
4	3340	77.3	1165	7	US-11-192-219-2	7
S	3318	76.8	868	7	US-11-166-730-3	m
ų	298.5	6.9	646	9	95	Sequence 695, App
7	298.5	. 6.9	708	9		'n
80	298.5	6.9	918	ø	US-10-995-561-696	69
o	250.5	5.8	836	9	US-10-821-234-1559 S	
10	210.5	4.9	979	9	US-10-636-320-6	
11	204	4.7	329	9	US-10-995-561-694	694
12	151.5	3.5	538	7	US-11-174-398-16	Sequence 16, Appl
13	149	3.4	2296	9	US-10-995-561-633	633
. 14	149	3.4	2355	9	US-10-995-561-623	623,
15	149	3.4	2355	9	US-10-995-561-627	627,
16	149	3.4	2384	ø	2	1545,
17	149	3.4	2386	9	US-10-995-561-626	626, A
18	145	3.4	1907	7	US-11-000-463-250 S	250,
19	144	3.3	384	7	US-11-075-351-32	32, A
70	141	9.3	1897	9	US-10-821-234-1635 S	
21	139.5	3.2	1433	7	40	Sequence 40, Appl
22	135.5	3.1	420	7	US-11-185-230-5	
23	135	3.1	313	۲	US-11-193-512-106	ĭ
24	132.5	3.1	2214	7		94
25	129.5	3.0	1315	9	US-10-995-561-630	630

ö

Gaps

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25; DB 7;

Score 4031; DB Pred. No. 0; 2; Mismatches

Query Match
Best Local Similarity 96.6%;
Matches 756; Conservative

US-11-192-219-7

2

Length 783; IndelB

Sequence 625, App Sequence 621, App Sequence 61, App Sequence 943, App Sequence 942, App Sequence 62, App Sequence 62, Appl Sequence 62, Appl Sequence 62, Appl Sequence 1587, Ap Sequence 1587, Ap Sequence 624, Appl Sequence 624, Appl Sequence 95, Appl Sequence 36, Appl Sequence 679, Appl Sequence 679, Appl Sequence 679, Appl Sequence 579, Appl Sequence 579, Appl Sequence 579, Appl Sequence 579, Appl
US-10-995-561-625 US-10-995-561-628 US-10-451-375-4 US-10-995-561-943 US-10-995-561-942 US-10-995-561-942 US-10-995-561-942 US-11-37-465-62 US-11-186-284-26 US-11-186-284-1587 US-11-186-214-1120 US-10-821-24-1120 US-10-995-561-995 US-11-135-885-35 US-11-135-885-35 US-10-995-561-995 US-11-135-885-35 US-11-1995-561-979 US-11-1995-561-979
1259 6 11386 6 11391 6 11391 6 11391 6 11391 6 11250 7 11250 7 1250 7 1303 7 13
26 122.5 27 122.5 28 122.5 29 120.5 30 117 32 111.5 34 111.5 36 111.5 37 110.5 38 111.5 39 101.5 41 107.5 42 107.5 43 107.5 44 105.5

```
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION: USES FOR WEX LIGANDS
ITILE OF INVENTION: USES FOR WEX LIGANDS
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
COUNTRY: USA
ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRITATION DATA:
APPLICATION NUMBER: US/11/192,219
FILING DATE: 27-Jul-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/667,197
FILING DATE: 20-Jun-1996
APPLICATION NUMBER: 08/585005
FILING DATE: 08-Jan-96
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: P-40,378
REFERENCE/DOCKET NUMBER: P0986P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1994
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ 1D NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 amino acids
TYPE: Amino Acid
Sequence 7, Application US/11192219
Publication No. US20050272656A1
```

Appli Appli Appli Appli Appli Appli Appli Appli Appli Appli

Sequence Sequence Sequence

Sequence Sequence Sequence Sequence Sequence

Sequence (

Sequence

Sequence Sequence Sequence Sequence

Perfect score:

Sequence:

OM protein

Run on:

Scoring table:

Searched:

ug-08-783-734d-10.rapbm

```
Length 894;
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Carter, Paul J.
APPLICANT: Carter, Paul J.
APPLICANT: Chiang, Nancy Y.
APPLICANT: Kyung, Jin Kim
APPLICANT: Ratthews, William
APPLICANT: Rodrigues, Maria L.
TITLE OF INVENTION: WSX RECEPTOR AGONIST ANTIBODIES
NUMBER OF SEQUENCES: 51
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WILDALIN (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/779,457
US-11-026-133-11
US-10-095-915-87
US-11-026-133-10
US-11-026-133-10
US-11-026-133-9
US-11-026-133-9
US-11-026-133-8
US-11-026-133-8
US-11-026-133-8
US-11-026-133-8
US-10-095-929-3
US-10-037-624A-4
US-10-373-624A-4
US-10-373-624A-4
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
US-10-373-624A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98.8%; Score 4271;
                                                                                                                                                                                                                                                                                                  ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
CLASSIFICATION BATA:
APPLICATION DATA:
APPLICATION NUMBER: 08/667197
FILING DATE: 06/20/96
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585005
FILING DATE: 01/08/96
ATTORNEY/AGENT INFORMATION:
NAME: Lee, Wendy M.
REGISTRATION NUMBER: 40,378
REFERENCE/DOCKET NUMBER: P0986P2
TELEPHONE: 415/525-1994
TELEPHONE: 415/525-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                         Sequence 51, Application US/08779457 Publication No. US20020193571A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 910/371-7168
INPORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 894 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch
    11165
11167
11167
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11161
1134
11234
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11234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ,
US-08-779-457-51
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       3340
33333
33228
33328
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33228
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3223
3223
3223
3123
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       Sequence 51, Appl
Sequence 2, Appli
Sequence 12, Appl
Sequence 51, Appl
Sequence 12, Appl
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                                                                                                                            (without alignments)
4764.376 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                       US-08-783-734D-10
4321
1 MMCQKFYVVLLHWEFLYVIA......IPSNVKKFYIHGMCTVLFMD 805
                                                                                                             January 2, 2006, 13:45:24; Search time 70.5974 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Published Applications AA Main:*
1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
                          GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-779-457-51

US-10-079-625-2

US-10-095-929-12

US-10-095-929-12

US-10-079-625-43

US-10-079-625-43

US-10-026-579-2

US-08-779-457-7

US-10-226-579-7

US-10-221-710-7

US-09-116-676-10

US-09-116-676-10

US-09-116-676-10

US-10-214-802-3

US-10-214-802-3

US-10-214-802-3

US-10-214-802-3

US-10-214-802-3

US-10-214-802-3

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4

US-10-214-802-4
                                                                                                                                                                                                                                                                                                             Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                             1867569 seqs, 417829326 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                      - protein search, using sw model
                                                                                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                            Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query
Match Length DB
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4271 4271 4271 4271 4271 4271 4031 3355

Result

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Tue Jan 3 12:13:52 2006
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US-08-599-455B-2
3328
3328
3328
3328
 Appli
                                                                   (without alignments)
3121.466 Million cell updates/sec
                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                          2, 2006, 10:19:03 ; Search time 21.3214 Seconds
                                                                                                         1 MMCQKFYVVLLHWEFLYVIA.....IPSNVKKFYIHGMCTVLFMD 805
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2,
                                                                                                                                                                                                                                                                                                                                                                       Description
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1
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Sequence 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                  /cgn2 6/ptodata/1/iaa/5_COMB.pep:*
/cgn2 6/ptodata/1/iaa/6_COMB.pep:*
/cgn2 6/ptodata/1/iaa/H_COMB.pep:*
/cgn2 6/ptodata/1/iaa/PCTUS COMB.pep:*
/cgn2 6/ptodata/1/iaa/RE COMB.pep:*
/cgn2 6/ptodata/1/iaa/RE COMB.pep:*
          GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                    US-08-618-957A-12
US-09-137-132-2
US-09-084-410-2
US-08-081-23D-2
US-08-583-153A-2
US-08-583-153A-2
US-08-583-153A-2
US-08-59-929-12
US-08-599-455B-43
US-09-09-781B-43
US-09-09-781B-43
US-09-094-410-43
US-09-094-410-43
US-08-640-389A-12
US-08-640-389A-12
US-08-640-389A-12
US-08-6803-346-1
                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                 572060 seqs, 82675679 residues
                                                                                                                                                                                                                                                                                                                                               SUMMARIES
                                                                                                                                                                                                          Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                           - protein search, using sw model
                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                              Issued Patents AA:*
                                                                                                                                                                                 Minimum DB seq length: 0 /
Maximum DB seq length: 200000000
                                                                                           US-08-783-734D-10
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Match Length DB
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                                                                                                    Perfect score:
                                                                                                           Sequence:
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                                                           Run on:
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222
223
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### TTT GIVINE

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UG-699-455B-2

Sequence 2, Application US/085995B

Patent No. 597261

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Texper. Robert 1.

APPLICANT: Texper. Robert 1.

APPLICANT: Texper. Robert 1.

APPLICANT: Texper. Robert 2.

CORRESONDENCE 4.

COURTS: WAS COUNTY: Read-able FORM:

WEDIN TYPE: Diskette

COUNTY: READ-ABLE FORM:

WEDIN TYPE: Diskette

COUNTY: 22 - Franklin Street

COUNTY: BARENT APPLICATION DATA:

APPLICATION WHORE: US/08/599, 455B

FILING DATE: 12 - DEC-1995

APPLICATION NUMBER: 08/56, 622

FILING DATE: 11 - DEC-1995

APPLICATION NUMBER: 08/56, 622

FILING DATE: 11 - DEC-1995

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - DEC-1995

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - DEC-1995

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - DEC-1995

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - SAC-1995

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - SAC-1996

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - SAC-1996

APPLICATION NUMBER: 08/56, 633

FILING DATE: 11 - SAC-1996

ATCORDERAL INFORMATION: AMILE INFORMATION: TELEBERK: 617-542-8906

TELEBERK: 617-542-8906

TELEBERK: MAIN-ATTENTICE:

LERGISTRATION POR SEQ ID NO: 2:

SEQUENCE: TERESONE CHARACTERISTICS:

LENGTH: EMANCE CHARACTERISTICS:

LENGTH: TELEBER: MAIN-AGAIGA

TURES AMIN-O-MAIN-O-MAILE

LENGTH: MASS AMIN-O-MAILE

LENGTH: MASS AMIN-O-MAILE

LENGTH: BASA AMIN-O-MAICH

TELEBER: AMIN-O-MAIN-O-MAILE

LENGTH: BASA AMIN-O-MAICH

TELEBER: AMIN-O-MAIN-O-MAILE

LENGTH: BASA AMIN-O-MAICH

TOPOLOGY: LINEAR
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us-08-783-734d-84.rapbn

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January 2, 2006, 13:46:09; Search time 8:94526 Seconds (without alignments) 748.439 Million cell updates/sec
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                                                                                                                                                                                                                        1 MMCQKFYVVLLHWEFLYVIA........BKNCSWAQGLNFQKRTDTL 894
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/cgn2_6/ptodata/2/pubpaa/USI1_NEW_PUB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
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US-11-166-730-3
US-10-666-730-3
US-10-995-561-696
US-10-995-561-696
US-10-636-220-6
US-10-636-220-6
US-11-094-519A-40
US-11-094-519A-40
US-11-004-519A-40
US-11-009-5561-694
US-11-009-995-61-633
US-10-995-661-633
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                                                                                                                                                                                                                                                                                                                57103 segs, 7488799 residues
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Maximum Match 100%
Listing first 45 summaries
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seq length: 200000000
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Maximum DB E
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WinPatin (Genentech) CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/192,219
FILING DATE: 27-U11-2005
CLASSIPECATION 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/667,197
FILING DATE: 20-UNN-1996
APPLICATION NUMBER: 08/585005
FILING DATE: 08-Jan-96
ATTORNEY/AGENT INFORMATION: GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Matthews, William
TITLE OF INVENTION: USES FOR WSX LIGANDS
NUMBER OF SEQUENCES: 47
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STREE: California
COUNTRY: USA REFERENCE/DOCKET NUMBER: P0986P1 TOPOLOGY: Linear

SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-11-192-219-7 NAME: Lee, Wendy M. REGISTRATION NUMBER: P-40,378 TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-1994 LENGTH: 783 amino acide TYPE: Amino Acid Sequence 7, Application US/11192219 Publication No. US20050272656A1 TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS: US-11-192-219-7

Gaps

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Score 4031; DB 7; Length 783; Pred. No. 0; 2; Mismatches 25; Indels

Query Match
Best Local Similarity 96.6%;
Matches 756; Conservative

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GenCore version 5.1.6

Upper Copyright (c) 1993 - 2006 Compugen Ltd.

Weeking Evency Compugen Ltd.

Second Compugen Ltd.

Second Copyright (c) 1993 - 2006 Compugen Coll Updates/second Copyright (c) 1994 Copyright (c) 1995 Copyright (c) 1994 Copyright (c) 1995 Copyright (c) 1994 Copyright (c) 1995 Copyrigh
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Searched: / 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 186756

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Listing first 45 summaries

Published Applications AA\_Main:\*
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2. /cgn2\_6/ptodata/1/pubpaa/US08\_PUBCOMB.ppp:\*
3. /cgn2\_6/ptodata/1/pubpaa/US08\_PUBCOMB.ppp:\*
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5. /cgn2\_6/ptodata/1/pubpaa/US10B\_PUBCOMB.ppp:\*
6. /cgn2\_6/ptodata/1/pubpaa/US10B\_PUBCOMB.ppp:\*
6. /cgn2\_6/ptodata/1/pubpaa/US10B\_PUBCOMB.ppp:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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,	Description	Sequence 51, Appl	Sequence 2, Appli	51,	12	12,	43,	2,	7	Sequence 7, Appli	Sequence 7, Appli	'n	Sequence 3, Appli	'n	10	3, 1	10,	10,	4,	Sequence 4, Appli	4,	7	Sequence 1, Appli	1	ñ	4,	•••	Sequence 73, Appl
SUMMARIES	ID	US-08-779-457-51	US-10-079-625-2	US-10-921-710-51	US-10-095-929-12	US-11-026-133-12	US-10-079-625-43	US-10-226-579-2	US-08-779-457-7	US-10-214-802-7	US-10-921-710-7	US-08-779-457-3	US-10-214-802-3	US-10-373-624A-2	US-10-774-721-10	US-10-921-710-3	US-10-095-929-10	US-11-026-133-10	US-08-779-457-4	US-10-214-802-4	US-10-921-710-4	US-08-779-457-2	US-09-894-039-1	US-10-095-929-11	US-10-214-802-2	US-10-226-579-4	US-10-921-710-2	US-10-893-315-73
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de	Query Match ]	100.0	100.0	100.0	6.66	99.9	99.5	99.4	84.1	84.1	84.1	79.4	79.4	79.4	79.4	79.4	79.1	79.1	79.0	79.0	79.0	79.0	79.0	79.0	79.0	79.0	79.0	79.0
	Score	4793	4793	4793	4788	4788	4768	4762	4031	4031	4031	3804	3804	3804	3804	3804	3792	3792	3785	3785	3785	3785	3785	3785	3785	3785	3785	3785
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6	2.0	78.8	78.8	78.8	78.8	78.7	78.7	78.7	78.6	78.6	76.9	76.9	76.9	76.9	76.9	76.2	73.4	8.69	
	3/85	3778	3775	3775	3775	3773	3773	3771	3768	3768	3687	3687	3687	3687	3687	3653	3517.5	3345	
6	P	53	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	

### LIGNMEN

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Sequence 51, Application US/08779457

Sequence 51, Application US/08779457

Publication No. US20020193571A1

GENERAL INFORMATION:

APPLICANT: Carter, Paul J.

APPLICANT: Carter, Paul J.

APPLICANT: Matthews, William

APPLICANT: Matthews, William

APPLICANT: Matthews, William

APPLICANT: Matthews, William

APPLICANT: Matthews, Malliam

APPLICANT: Matthews, Milliam

APPLICANT: South San Exceptor AGONIST ANTIBODIES

CORRESPONDENCE ADDRESS:

ADDRESSEE: Genentech, Inc.

STREET: AGO Point San Bruno Blvd

CITY: South San Francisco

STREET: AGO Point San Bruno Blvd

CITY: South San Francisco

STREET: AGO Point San Bruno Blvd

CONFUTER READBLE FORM:

MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMFUTER: IBM PC compatible

COMFUTER: IBM PC compatible

COMFUTER: IBM PC compatible

COMFUTER: Minpatin (Genentech)

COMFUTER: Minpatin (Genentech)

CLASSIFICATION NUMBER: US/08/779,457

FILING DATE:

CLASSIFICATION NUMBER: 08/667197

FILING DATE:

APPLICATION NUMBER: 08/667197

FILING DATE: 01/08/96

ATTORNEY/AGENT INFORMATION:

MANE: Lee, Wendy N.

RESISCONTANICATION INFORMATION:

TELECOMMUNICATION OF SI:

SEQUENCE SISCONERY LINEAR:

SEGUENCE SISCONERY LINEAR:

SERGISTER AMINO Acid

TTELET AMINO Acid

TTELET AMINO Acid

TTELET AMINO Acid

TOPOLOGY: Linear

US-08-779-45-51
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Query Match

100.0%; Score 4793; DB 2; Length 894;

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GenCore version 5.1.6
Copyright (c) 1993 - 2006 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                          OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Match Length
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                                                                                                                                                      Title:
Perfect score:
                                                                                                                                                                                                                    Scoring table:
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                                                                                                                                                                                        Sequence:
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                                                                                           Run on:
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No.
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28	3792	79.1	896	N	US-09-357-914-33	Sequence	33,	App]
29	3792	79.1	968	~	US-10-095-929-10	Sequence	, 0,	Appl
30	3792	79.1	868	н	US-08-693-697-36	Sequence	36,	Appl
31	3785	79.0	923	~	US-08-780-562-4	Sequence	4	Appli
32	3785	79.0	1165	ч	US-08-599-455B-4	Sequence	4,	Appli
33	3785	79.0	1165	~	US-09-093-814-1	Sequence	'n	Appli
34	3785	79.0	1165	~	US-09-069-781B-4	Sequence	4,	Appli
35	3785	79.0	1165	~	US-08-618-957A-11	Sequence	11,	Appl
36	3785	79.0	1165	7	US-09-137-132-4	Sequence	4	Appli
37	3785	79.0	1165	~	US-09-094-410-4	Sequence	4,	App1:
38	3785	79.0	1165	N	US-08-708-123D-4	Sequence	4,	Appli
39	3785	79.0	1165	~	US-08-583-153A-4	Sequence	4	Appli
40	3785	79.0	1165	8	US-08-570-142D-4	Sequence	4	Appli
41	3785	79.0	1165	N	US-08-780-562-2	Sequence	7	Appli
42	3785	79.0	1165	~	US-08-638-524B-4	Sequence	4,	App1
43	3785	79.0	1165	~	US-10-095-929-11	Sequence	ij	App]
44	3779	78.8	896	Н	US-08-640-389A-10	Sequence	10,	App]
45	3775	78.8	898	N	US-08-588-189-3	Sequence	e,	Appli

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Sequence 2, Application US/0859455B

Sequence 2, Application US/0859455B

Sequence 2, Application US/0859455B

Patent NO. 597561

GENERAL INFORMATION

PEDLICANT: Tartsqlia, Louis A.

APPLICANT: Clipepper, Manice A.

ITIELS OF INVENTION: MODULATE BODY WEIGHT USING THAT

ITIELS OF INVENTION: MODULATE BODY WEIGHT USING THAT

ITIELS OF INVENTION: MODULATE BODY WEIGHT USING THE OB RECEPTOR

NUMBERS OF SEQUENCES: 44

COUNTRY: US

COUNTRY: US

COUNTRY: US

COUNTRY: US

COUNTRY: US

COUNTRY: US

COUNTRY: IS Diskette

COMPUTER: IS IN COMPATION BOTH

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APPLICATION NUMBER: 08/56,663

APPLICATION NUMBER: 08/56,663

FILING DATE: 11-DEC-1995

APPLICATION NUMBER: 08/56,663

FILING DATE: 08/56,010

FILENCENTE: 08/56,01
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